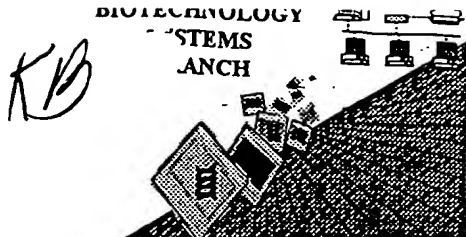


PCT09

## CRF Problem Report



The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: 09/937,204A  
Filing Date: 09/21/2001  
Date Processed by STIC: 11/25/02

STIC Contact: Mark Spencer, 703-308-4212

### Nature of Problem:

The CRF (was):

- ☒ (circle one) Damaged or Unreadable (for Unreadable, see attached)  
☐ Blank (no files on CRF) (see attached)  
☐ Empty file (filename present, but no bytes in file) (see attached)  
☐ Virus-infected. Virus name: \_\_\_\_\_ The STIC will not process the CRF.  
☐ Not saved in ASCII text  
☐ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.  
☐ Did not contain a Sequence Listing. (see attached sample)  
☐ Other: \_\_\_\_\_

PLEASE USE THE CHECKER VERSION 3.1 PROGRAM TO REDUCE ERRORS.  
SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name,  
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,  
2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service , or other delivery service to: U.S. Patent and Trademark Office,  
B x Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



**Does Not Comply**  
**Corrected Diskette Needed**

PCT09

**RAW SEQUENCE LISTING**

DATE: 04/25/2002

PATENT APPLICATION: US/09/937,204

TIME: 11:37:43

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt

Output Set: N:\CRF3\04252002\I937204.raw

4 <110> APPLICANT: Fritig, Bernard  
 5 Toquin, Valerie  
 6 Geoffroy, Pierrette  
 7 Legrand, Michel  
 8 Kauffmann, Serge  
 11 <120> TITLE OF INVENTION: INDUCIBLE COMTII PROMOTER, CHIMERA GENE  
 12 CONTAINING SAME AND TRANSFORMED PLANTS  
 15 <130> FILE REFERENCE: A34638-PCT-USA  
 17 <140> CURRENT APPLICATION NUMBER: US 09/937,204  
 18 <141> CURRENT FILING DATE: 2000-03-22  
 20 <150> PRIOR APPLICATION NUMBER: PCT/FR99/03700  
 21 <151> PRIOR FILING DATE: 1999-03-22  
 23 <150> PRIOR APPLICATION NUMBER: PCT/FR99/07646  
 24 <151> PRIOR FILING DATE: 1999-06-11  
 26 <160> NUMBER OF SEQ ID NOS: 26  
 28 <170> SOFTWARE: FastSEQ for Windows Version 4.0

**ERRORED SEQUENCES**

712 <210> SEQ ID NO: 26  
 713 <211> LENGTH: 30  
 714 <212> TYPE: DNA  
 715 <213> ORGANISM: Artificial Sequence  
 717 <220> FEATURE:  
 718 <223> OTHER INFORMATION: Synthetic Oligonucleotide PS14  
 720 <400> SEQUENCE: 26  
 721 acgcgtcgac cagtggtag tttagctgtc 30  
 E--> 734 ny02:358522.1 -delete

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/09/937,204

DATE: 04/25/2002

TIME: 11:37:44

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt

Output Set: N:\CRF3\04252002\I937204.raw

Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of <220> to <223> is MANDATORY if n's or Xaa's are present.

in <220> to <223> section, please explain location of n or Xaa, and which  
residue n or Xaa represents.

Seq#:26; N Pos. 31

## VERIFICATION SUMMARY

DATE: 04/25/2002

PATENT APPLICATION: US/09/937,204

TIME: 11:37:44

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt

Output Set: N:\CRF3\04252002\I937204.raw

L:44 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:47 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:51 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:59 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:63 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:67 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:71 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:75 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:79 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:82 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:86 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:89 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:92 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:96 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1  
L:143 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2  
L:146 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2  
L:149 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2  
L:152 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2  
L:447 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:12  
L:536 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14  
L:734 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:26  
L:734 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:26  
L:734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:30  
L:734 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:34 SEQ:26  
L:734 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10  
L:734 M:112 C: (48) String data converted to lower case,  
L:734 M:252 E: No. of Seq. differs, <211> LENGTH:Input:30 Found:34 SEQ:26